



52288B-SEQ-listing.txt
SEQUENCE LISTING

<110> Hillyard, Jeanna
Roberts, James
Ye, Minwei

<120> Cotton Event PV-GHBK04 (757) and Compositions and Methods for Detection Thereof

<130> 38-21 (52288)B

<140> 09/990,659

<141> 2001-11-16

<150> US60/249,757

<151> 2000-11-17

<160> 21

<170> PatentIn version 3.0

<210> 1

<211> 20

<212> DNA

<213> artificial sequence

<220>

<221> misc_feature

<222> (1)..(20)

<223> a 5' genome-insert junction nucleotide sequence which is/is complementary to a sequence diagnostic for nucleic acids derived from the cotton event 757 recombinant genome

<400> 1

gtttgcttgg acactgatag

20

<210> 2

<211> 20

<212> DNA

<213> artificial sequence

<220>

<221> misc_feature

<222> (1)..(20)

<223> a 3' genome-insert junction nucleotide sequence which is/is complementary to a sequence diagnostic for nucleic acids derived from the cotton event 757 recombinant genome

<400> 2

aaacccttcc tggaaaaata

20

<210> 3

<211> 20

<212> DNA

<213> Gossypium hirsutum

<220>

<221> misc_feature

<222> (1)..(20)

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<400> 3
tgttctgtgg aaaaggaagg

20

<210> 4
<211> 20
<212> DNA
<213> Gossypium hirsutum

<220>
<221> misc_feature
<222> (1)..(20)

<400> 4
atgcctgcag gtcaattcaa

20

<210> 5
<211> 138
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(138)
<223> part of a 5' non-functional sequence inserted into the cotton genome in cotton event 757

<400> 5
acactgatag tttaaactga aggcgggaaa cgacaatctg atcccagctt gcatgcctgc 60
aggtaattc aatattgtgg caggacattg ctacatgata cctcttagaa ttgttagac 120
ttcagatcga tcttgtca 138

<210> 6
<211> 767
<212> DNA
<213> Gossypium hirsutum

<220>
<221> Unsure
<222> (1)..(767)
<223> 5' cotton (Gossypium hirsutum) genome sequence

<400> 6
gtccgggggg cttatcctgt attcatttgc acccacataa acagccaaat taaccaaacc 60
catattcaac tgaaactccc aaagccattc ctactttagc ttttcaccca ctaactcaaa 120
agaaaaacact cacctagctt cttgcttt tcttttggat tgtttagat ctacaaaaag 180
atgattcaag aactccttgg aggttcttct tgcttaact ttggagggga gaggaagatc 240
tccatcaatg gaagcatttt ggaaggaacc cccacttctt ctccatcacc atcatttct 300
tcttcttcgg cgacgacttc atcgaccact aattcatcga atccggagaa tcattaccag 360
aatttgaggt gccccaggtg tgattcctcc aacacaaagt tctgctatta caacaactac 420
aacctcactc agcctcgtca ctttgcaag acttgccgtc ggtattggac caaaggagga 480

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gctctcagaa acgttcctat tgggtgggg tgtagaaaa acaaaaagcac tactggtgtt	540
tcaacatctc tggggaaatc aacttcttcc aagataaaa cagtagttc tgaaattgga	600
agatctgggt tcgatcatga gcttcagtct actccaattc tttggacttc agcgcccag	660
acttccatc ttctatccaa tctaacctca atgagagcta ccctaaaccc taaccctaac	720
acattgtcta accctgttag tattaaggaa gaagtgagtt tgcttgg	767

<210> 7	
<211> 206	
<212> DNA	
<213> artificial sequence	
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<221> misc_feature	
<222> (1)..(206)	
<223> part of arbitrarily assigned 3' end DNA sequence inserted into the cotton event 757 genome.	

<400> 7	
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tctttttgga atgctgctcc gtcgtcaggc tttccgacgt ttgggtgggtt gaacagaagt	120
cattatcgca cggaatgccca agcaactcccg agggaaaccc tgtggttggc atgcacatac	180
aatggacga acggataaac ctttc	206

<210> 8	
<211> 307	
<212> DNA	
<213> Gossypium hirsutum	
<220>	
<221> Unsure	
<222> (1)..(307)	
<223> 3' cotton (Gossypium hirsutum) genome sequence	

<400> 8	
tggaaaaata atcaacacca cgctcaacaa caacagaata ataatgggtt cttttaggt	60
gaagttcaaa acacaggtat tcaagaactg tatcaaaggc tcaaatcatc atcaagttat	120
tactctgata ctccagcagt aattctaagc aatgtcgctt cttcttcatc aacatccatt	180
ttggagtcag ctccagttgc tggggagaa ttgggttaact ggaatccggc atttcatca	240
tcgtggtctg atcttccaac aactaatggt gcatacctt aaaataaccc tttacccccc	300
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<210> 9	
<211> 26	
<212> DNA	
<213> artificial sequence	

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<220>
<221> misc_feature
<222> (1)..(26)
<223> 5' cotton (*Gossypium hirsutum*) genome PCR primer

<400> 9
gagagagata ggcactaaag taagca

26

<210> 10
<211> 28
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(28)
<223> 5' insert PCR primer

<400> 10
ttagacaaat tgtcacggtc taccagaa

28

<210> 11
<211> 24
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(24)
<223> 3' insert PCR primer

<400> 11
ttcccaacga tcaaggcgag ttac

24

<210> 12
<211> 27
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(27)
<223> 3' cotton (*Gossypium hirsutum*) genome PCR primer

<400> 12
ttgatgcact tacgaaaagaa gaaccga

27

<210> 13
<211> 905
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(905)
<223> 5' cotton (*Gossypium hirsutum*) genome + insert sequence

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agaaaacact cacctagctt ctttgctttt tcttttggat tgtttagat ctacaaaaag 180
atgattcaag aactccttgg aggttcttct tgcttaaact ttggagggga gaggaagatc 240
tccatcaatg gaagcatttt ggaaggaacc cccacttctt ctccatcacc atcatcttct 300
tcttcttcgg cgacgacttc atcgaccact aattcatcga atccggagaa tcataaccag 360
aatttgaggt gccccaggtg tgattcctcc aacacaaaagt tctgctatta caacaactac 420
aacctcactc agcctcgta ctttgcaag acttgcgtc ggtattggac caaaggagga 480
gctctcagaa acgttcctat tgggtgggg tgttagaaaa acaaaaagcac tactgggttt 540
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agatctgggt tcgatcatga gcttcagtct actccaattc tttggacttc agcggcccag 660
acttcccatc ttctatccaa tctaacctca atgagagcta ccctaaaccc taaccctaac 720
acattgtcta accctgttag tattaaggaa gaagttagtt tgcttggaca ctgatagttt 780
aaactgaagg cgggaaacga caatctgatc ccagcttgca tgcctgcagg tcaattcaat 840
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tgtca 905

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<210> 14
<211> 513
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(513)
<223> 3' cotton (Gossypium hirsutum) genome + insert sequence

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<400> 14
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cattatcgca cggaaatgcca agcaactcccg agggaaaccc tgggttggc atgcacatac 180
aaatggacga acggataaac ctttctgga aaaataatca acaccacgct caacaacaac 240
agaataataa tgggttcctt gtaggtgaag ttcaaaaacac agttattcaa gaactgtatc 300
aaaggctcaa atcatcatca agttattact ctgataacttc agcagtaatt ctaagcaatg 360
tcgcttcttc ttcatcaaca tccatTTGG agtcagctcc agttgctggg ggagaattgg 420
gttactggaa tccggcattt tcattatcgt ggtctgatct tccaaacaact aatggtgcat 480

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atccttaaaa taacccttta cctttcggtt aat

513

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<210> 15
<211> 4973
<212> DNA
<213> artificial sequence

.
<220>
<221> misc_feature
<222> (1)..(4973)
.
<223> sequence of 5' flank to full-length cry1Ac coding region
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<400> 15
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accctaacac attgtctaac cctgttagta ttaaggaaga agtgagttt cttggacact
gatagttaa actgaaggcg ggaaacgaca atctgatccc agcttgcatt cctgcaggc 120
aattcaatat tgtggcagga cattgctaca tgataccctct tagaattgtt tagacttcg 180
atcgatcttgc tcaagtctgaa agacccaaaa acaaattgcaa tttctttctt ggttagaccgt
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1380
1440
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ggtcagtaca tcccacatcg atgtccaagg agaagtgtat agaatgggtgg gcacacttct	1740
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<210> 16
<211> 19
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(19)
<223> 5' primer to 5' flanking sequence of SEQ ID NO: 15 from 8 to 26;

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<400> 16
gacttccat cttctatcc

19

<210> 17
<211> 19
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(19)
<223> 3' primer to partial e35S promoter of SEQ ID NO: 15 from 3154 to 3
136

<400> 17
attgtgcgtc atcccttac

19

<210> 18
<211> 22
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(22)
<223> 5' primer to partial 3' cry1Ac sequence of SEQ ID NO: 15 from 2581
to 2603

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22

<210> 19
<211> 20
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(20)
<223> 3' primer to partial 5' cry1Ac sequence of SEQ ID NO: 15 from 3455
to 3435

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<210> 20
<211> 19
<212> DNA
<213> artificial sequence

<220>
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<400> 20
ataaaaggaaa ggccatcgt 19

<210> 21
<211> 25
<212> DNA
<213> artificial sequence

<220>
<221> misc_feature
<222> (1)..(25)
<223> 3' primer to full-length cry1Ac sequence of SEQ ID NO: 15 from 497
3 to 4949

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10